SEQUENCE LISTING

52
100
148
196
244

tta tct ttg att gtc ttg agt gct gca tca gct aag aca aaa acc aca Leu Ser Leu Ile Val Leu Ser Ala Ala Ser Ala Lys Thr Lys Thr Thr 80 85 90

gag tca gag gga aaa aaa acg tcc tgatgaggat tgtgcaattt ccggaccatc 346 Glu Ser Glu Gly Lys Lys Thr Ser

406

466 526

586

646 706

766

826

886

946

1006

1066

1126 1186

1246 1306

1366

1426

1486 1546

1606 1666

1726

1786

1846

1906

1966

2026 2086

2146

2206

2266 2326

2334

95

attttttaaa aattataaat tatgaaatcc cacattttca atcccaattt ctggaacgtg ttttattttg agcacagaat ggcaacatcc caggaaaaaa agtcatgctc ccattttgct tqtaatcaaq tqaqctggaa ctgaccctac cccaaatatt ttttgaatag ggaaaagact caactggacc cetetaagga etgggagetg geatggaget ggeatettet gagactgaet tgagaagagc ctgataacgc ctagaggaaa caggcagggt tttgcgagca ggggaagatg ataqtqqqtq qqtqqqqaqc tqqcqaqqqt gccccaggca gaggcaccgt gtgtgtgcaa aggcctgcag gtggagaagg gcctgggact cttggagaat ggcaggaagt ttggtgtgcc tgtagtctat gagccaggct cagggcagca aaggtctgtc ctgcaggtgt tgtgatgagc tqtaccactt agtgggcacc atcaagatga acagagagta acacggtggc actgagaact tgagaacage teactetaga atgaactgtg teeteeaaag tgtgeagage caatacetag gggtccccaa ggtgactgag cacgggcaca gatccagcag caaatccccc cagtccaaga getgttettt ceattetetg ttettteeat tetetgttet tetggteett etgettatgg caaggtgaaa gtcacaggtg gaattgtccc tatcacctct cccacaccct gatctccttt tacaacaaag agcaagcatc ctctacaaca aagcetttgg ttggtgtcag tgcctggctg ggaggaagta actgttgttt ttactgtgtt taatttcact cctgccgtct gttcacqqca ccaqtqatca qqttctctqc caqtqqqaqt gatagaaagt taccttttta aagtaaattt cttggaacgc aaaaaacaag ccaagttaaa taaaaataca aaatatgggg ccaggcgcgg tggctcgtgc ctgtcatccc agcactttgg gaggctgaga cggtggatca cctgaggtca qqaqtttqaq accaqcctqa ccaacaaqqt qaaqccccqt ctctactaaa aatacaaaaa ttagccgggc gtggtggcag gcacctgtag tcccagctac tcgggaggct gggacagagg aattqcttqa acccqqqagg cggaggttgc agtgagccga gatcacgcca ccactgcact ccagcctggg tgacggagcg agatgccatt tcaaaacaaa aacaaaatat gtactggtac cagtacacag taggaaggtg ggcaaaactt gggaaggggg atattcaaag gacagggttt gggaaatgct ggatcaaggt cggggaagaa ggagaactga gaggctgtta taatttagag aagtgcttct cagagtgggg gccagcagcc aggcgccgtg gctcatgcct gtaaccttaa cactttggga ggtctaggcg ggaggattgc ctgagcccag gagttcgagt ccagcttgtg caacatagtg agatgctgtc tctacaaaaa atttaaaaat tagctggtgt cctctcagtg tgtcttgtcc tctccatgtt tctaaaataa aggaagaaag gcccagcgca gtggcgtaca cctatagtct cagcactttg ggaggccaag gtgggcagat cacttgaggt caggagttcg agaccagcct ggctaacatg gcaaaacctt gtttctactg gaaatacaaa aattagctag qcqtqqtgqt gcacgcctgt aatcccagct acttgggagg ctgagggagg agaaccgctt qaqcctqqqa qqcaqaqqct qcaqtqaqcc aagatcacac actgcactcc agcctgggtg acagagogag actocatoto aaataaataa ataaataaat aaaaataaat acataaatac ataaaata

<210> 2 <211> 99 <212> PRT <213> Homo sapiens <400> 2 Met Cys Cys Trp Pro Ser Pro Trp Val Gln Gly Ser Pro Gly Ile Trp 15 His Leu Trp Ala Val Leu Ala Cys His Leu Gly His Ser Ser Ser Arg Gln Gly Ile Leu Arg His Arg Pro Gly Gly Ala Leu Pro Ser Thr Pro 40 Gly Cys Thr Met Thr Ser Thr Leu Gly Gln Arg Pro Leu Leu Gln Gly 50 55 Cys Glu Asp Ile Met Val Gln Pro Glu Gly Asp Leu Ser Leu Ile Val 70 75 Leu Ser Ala Ala Ser Ala Lys Thr Lys Thr Thr Glu Ser Glu Gly Lys 85 90 Lys Thr Ser <210> 3 <211> 297 <212> DNA <213> Arificial Sequence <220> <223> This degenerate sequence encodes the amino acid sequence of SEQ ID NO:2. <221> variation <222> (1)...(297) <223> N is any nucleotide. <400> 3

atgtgytgyt ggccnwsnoc ntgggtncar ggnwsnocng gnathtggca yytntgggcn gtnytngcnt gycayytngg ncaywsnwsn wsnmgncarg gnathytnmg ncaymgnocn ggnggngcny tnocrwsnac nccnggntgy acnatgacnw snacnytngg ncarmgnocn ytnytncarg gntgygarga yathatggtn carcongarg gngayytnws nytnathgtn ytnwsngcng cnwsngcnaa racnaaracn acngarwsng arggnaaraa racnwsn 60 120

180 240

297

<210> 4 <211> 23

<212> DNA <213> Artificial Sequence	
<220> <223> PCR primer	
<400> 4 ctgatgcagc actcaagaca atc	23
<210> 5 <211> 24 <212> DNA <213> Artificial Sequence	
<220> <223> PCR primer	
<400> 5 ggcatttgtg ggcagtgttg gggc	24
<210> 6 <211> 18 <212> DNA <213> Artificial Sequence	
<220> <223> PCR primer	
<400> 6 ggagctggca tcttctga	18
<210> 7 <211> 18 <212> DNA <213> Artificial Sequence	
<220> <223> PCR primer	
<400> 7 tccccaccca cccactat	18
<210> 8	

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide linker

<400> 8

Gly Gly Ser Gly Gly Gly Gly Gly Ser Gly Gly Gly Ser 1 5 10 15